

OM nucleic - nucleic search, using sw model

Run on: June 2, 2003, 00:51:28 ; Search time 4674.33 Seconds  
(without alignments)  
16997.243 Million cell updates/sec

Title: US-09-596-194-59  
Perfect score: 2730  
Sequence: 1 gtcgacccacgcgtccgtcc.....aaaaaaaaaagggcgccgc 2730

Scoring table: OLIGO\_NUC  
Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 400

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:\*  
1: gb\_ba:\*  
2: gb\_htg:\*  
3: gb\_in:\*  
4: gb\_om:\*  
5: gb\_ov:\*  
6: gb\_pat:\*  
7: gb\_ph:\*  
8: gb\_pl:\*  
9: gb\_pr:\*  
10: gb\_ro:\*  
11: gb\_sts:\*  
12: gb\_sy:\*  
13: gb\_un:\*  
14: gb\_vi:\*  
15: em\_ba:\*  
16: em\_fun:\*  
17: em\_hum:\*  
18: em\_in:\*  
19: em\_mu:\*  
20: em\_om:\*  
21: em\_or:\*  
22: em\_ov:\*  
23: em\_pat:\*  
24: em\_ph:\*  
25: em\_pl:\*  
26: em\_ro:\*  
27: em\_sts:\*

search repeated in  
GenSeq  
US Pats  
US P&PUBS  
EST

GenCore version 5.1.4\_p5\_4578  
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Run on: June 2, 2003, 00:51:28 ; Search time 3446.67 Seconds  
(without alignments)  
16997.243 Million cell updates/sec

Title: US-09-596-194-60  
Perfect score: 2013  
Sequence: 1 atggcccagctgttcctgcc.....ccctgcagctctgggtcacc 2013

Scoring table: OLIGO\_NUC  
Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 400

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:\*  
1: gb\_ba:\*  
2: gb\_htg:\*  
3: gb\_in:\*  
4: gb\_om:\*  
5: gb\_ov:\*  
6: gb\_pat:\*  
7: gb\_ph:\*  
8: gb\_pl:\*  
9: gb\_pr:\*  
10: gb\_ro:\*  
11: gb\_sts:\*  
12: gb\_sy:\*  
13: gb\_un:\*  
14: gb\_vi:\*  
15: em\_ba:\*  
16: em\_fun:\*  
17: em\_hum:\*  
18: em\_in:\*  
19: em\_mu:\*  
20: em\_om:\*  
21: em\_or:\*  
22: em\_ov:\*  
23: em\_pat:\*  
24: em\_ph:\*  
25: em\_pl:\*  
26: em\_ro:\*  
27: em\_sts:\*

Search repeated in  
GenSeq  
US Pats  
US PGPubs  
EST